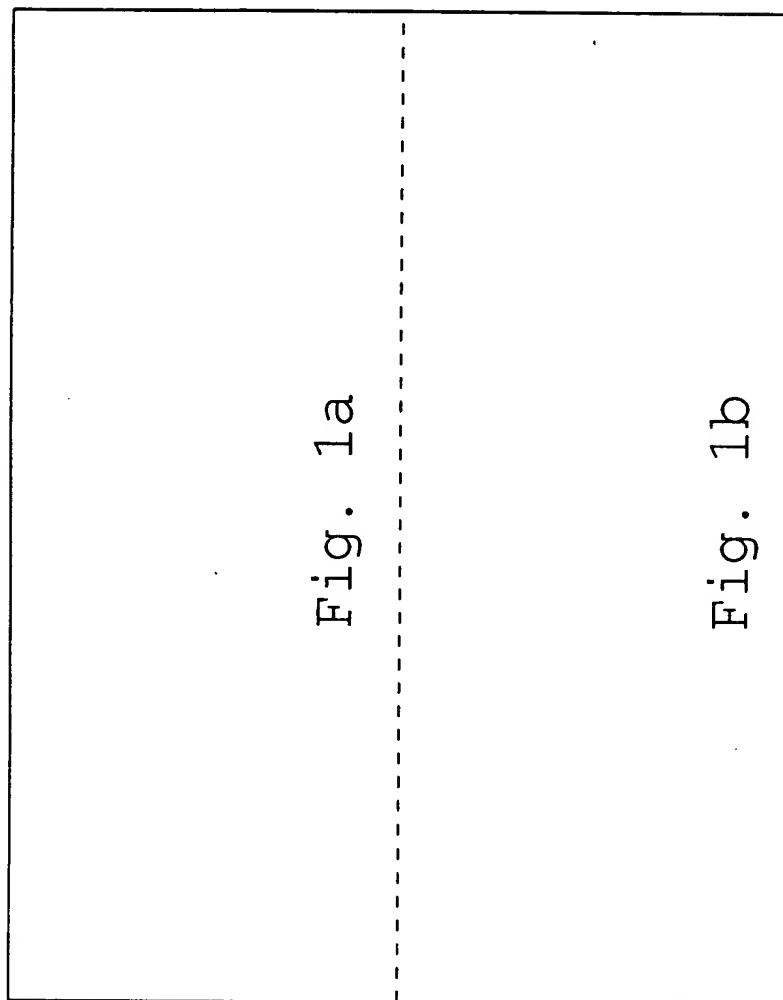


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106TDT DEEDOT



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TOGTFDT "DEEDOOT" (S)

SEQ ID NO:

6 Streptopain	1 MNKKKLGRLLSLLALGGLVLANPVFAQONFARNEKEAKDSATFIQKSAALKAARSAAE
5 prtT	1 --MKRIFVTLCLLLCLPML-QAGPVTRSKAEQTAKNFFAKROPTL---SSSTASURMDP
1 Periodontain	1 --MKKSELAAVMLFCTAMQGHSAVPTKERRALSLAR--LALRQVSLRMGQTAWSDKLSIDY
61 Streptopain	61 DIKLDKVNLG---GELSGSNMYYNISTG-GFVIVSGDKRSPEILGYSTSGSFDANGKE
55 prtT	55 VYK--AAER-----EE--ALFVFNRRGEKCGFLLVAADDRFPPEVIGYAFKGFHDAARIP
58 Periodontain	58 VYRQGDARIGITSQEEGSPAYFYVANRGNNEGVALVAADDRIPITLLAYSPIGRFDMDSMP
116 Streptopain	116 N-----IASFMESYVEQIKENKKLDTTYAGTAEIKQPVVKSLLDSKGIHNNQG
105 prtT	105 DNLRGWLKGYEREMLAAMDGKAEPIDPIREAKPTRDPSSIAPIETGEHASDPILMDQG
118 Periodontain	118 DNLRMMLQIYDQEIQLISGKAQNLNHEILR---TEGVPAEVEVHALMDNCHFANDPMMRNQ
164 Streptopain	164 NPYNLLTPVIEKVKPGEQSFVGOHAATGCVATATAQIMKYHNPKNKGLDYTYTLSSNNP
165 prtT	165 YPENTLHPLLP-----SQQAAYTGCVATAMQIMRHYKWPEKASGEYDYDDMTGT
175 Periodontain	175 YPMNNKEPLLP-----NGNHAYTGCVATAAQIMRYHSWPLQGECSLDYHAGSLVG
224 Streptopain	224 YFNHPKNLFAAISTRQYNWNNTLPTYSGR-ESNVQKMAISELMADVGISVDMDYGPSSGS
216 prtT	216 HT-----HYSGTGGETYNWSKMPGNISVG-LSPEVKAISTEMRDVSFSVNMQFADFSGG
226 Periodontain	226 N-----WSGTFGEMYDWIINMPGNPDLDNLTQSQVDAYATLMRDVSASVSMSEYENGSG
283 Streptopain	283 AGSSRVQRALKENFCYNQSVHQINRSDLSKQDMWAEQIDKELSQNQPVVYQGVGVKGHAF
270 prtT	270 TESSIFVERALRETFYKKSRLRYIHRSLLPCKEMKDMIRKELAENRPVYAGADGSMGHAF
279 Periodontain	279 TYSVAVVVVGALRNFRYKRSLLQLHVRALYTSQEWIEMIRGELASGRPVYAGANNQSGHAF
343 Streptopain	343 VIDGAGRNFFHVNVNWGWGVSQGFRELDA LNPSALGTGGGAGGFNGYQSAVVGIKP----
330 prtT	330 VCDGYEPDGTFFHFNVNWGWGVSNGNFYLNLLNPGSLGTRAGDGGYSTDOEVLVIGIEPASNE
339 Periodontain	339 VCDGYASDGTFFHFNVNWGWGVSNGFVKKLTLLSPASLIGGEGIGCFNIYQEIITTGIEPAKTP

Fig. 1a

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prtt
Periodontain
390 VP-GIVDPPTITLYGQHNMSDEA-LDSVKIKNYSTYAGDVKLKYRLTLPNGTETTNPA
399 AEAGTDALPIALKDIEAEYKSESGLNMGYSIYNTGEEQSNLDLYRLNKADEGEVIEVKT

prtt
Periodontain
448 VIVPIVMEDIIGESTGNIATPCSQFAEGKNTISILYRTDGMADWKEKHHILMGLVNKITEV
459 SSINISWYG-YGEHPESFSLAPNQLSQGINITITLYRRTGTTEQWEPVRRHAQGGYVNSIKV

prtt
Periodontain
508 TMPAGDVAYSAD---ARIVLKDGSLSHDLKAYSDCKLSATVYNPGTTEFRSRVTFALRN
518 NTTDPNNVVTVTDNNECKLSIVPNSFVADLNSYEHSTLTVQFNSDSPDEIRTPVAFALST

prtt
Periodontain
565 --TEGRIFLGRHVEVHPGDEDEGEKVSLLITGLKARACQYMLVCTGDMESLMEDASWIE
578 GATADDVISLGWVMAEVPGSSN-YPVVWSKDVLTLSECDYTLWYRFSIN--NQKDEWKK

prtt
Periodontain
623 LASIEVAEHTSTHSSLLVASNPQIDLLTVHRANPETLPTEFSTNECGGATFSCKEIVVAIK
635 TGSVSVKTPTEYTHPLFEVGNQTSYTLDMAHNRVLPDFTEKKNLG-LPFNGEIVVFRQ

prtt
Periodontain
683 AFSET-FFQAKEEHMSLAQGETKVLSPETANSSLYTNAELFPDGYIYIVREQG-FWDP
694 TQSSGSLWAAQETVHKQGETFVYKPVVEG-----PIPDGSRATLHAFFVNGQQQ

prtt
Periodontain
741 KDLFGDYRYRTRRLTDLSSSDTAGKDVSTIVLYPNPAEDYVHVAIPPTYAGSTLR LFDLQ
745 IY LK GK RNYTVKIANGTAVEAIE--SSEETIRVFPNPARDYVEISAPCIPQETSL LFDLS

prtt
Periodontain
801 GRMQSTKTESADMRVDVERLPKGTIIVVEDMVGKLFIR-
803 GKIVMKNSLSACHGRMDVSR LPN GAVILKV DGYTTKINIVH

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Fig. 1b

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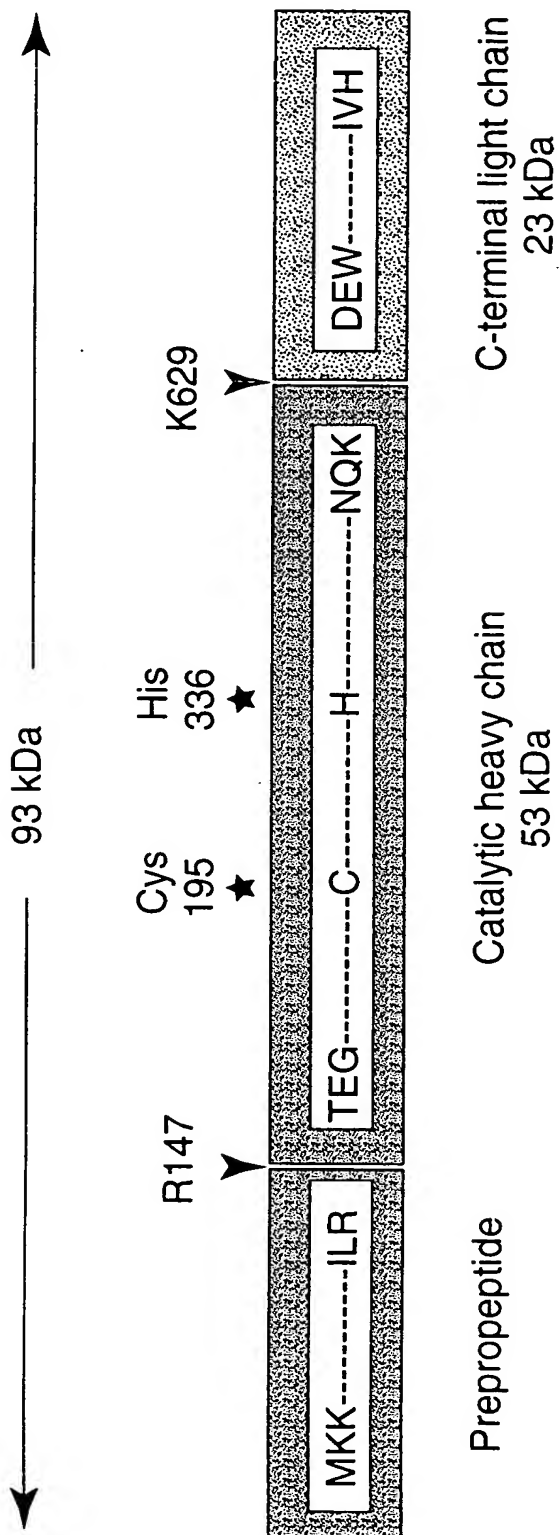


Fig. 2

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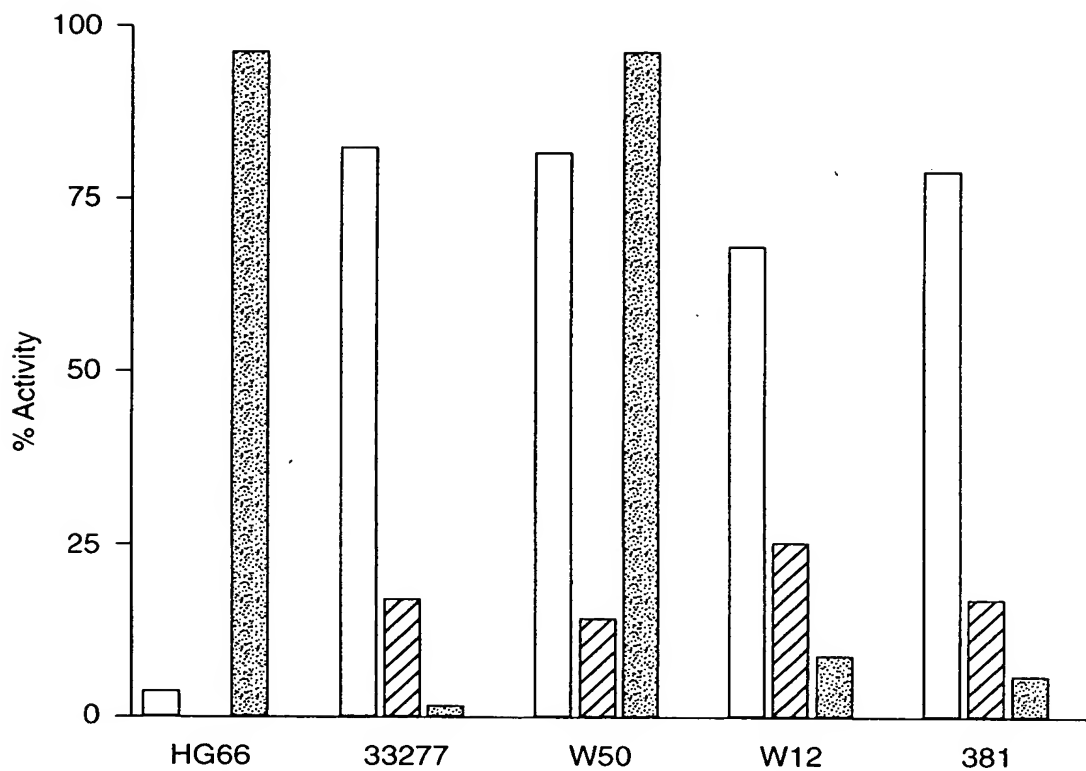


Fig. 3

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MPSSVSWGILLAGLCCLVPVSLAEDPQGDAQKTDTSHHDDQDHP
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YHSEAF TVNFGDTEEAKKQINDYVEKGTQGGKIVDLVKELDRDTVFALVNYIFFKGKWE
RPFVVDTEEEEDFHVQVTTVKVPMKRLGMFNIQHCKKLSSWVLLMKYLG NATAIFF
LPDEGKLQHLENELTHDIIITKFLNEDRRSASLHLPKLSITGTDLKSVLGQLGITKV
FSNGADLSGVTEEAPLKLSKAVHKAVLTIDEKGTEAAGAMFLEAIPMSIPPEVKFNKP
FVFLMIEQNTKSPLFMGKVVNPTQK

SEQ ID NO:3

Fig. 4

GAMFLEAIPMSIPPE

SEQ ID NO:4

Fig. 6

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Fig. 5

ATGAAAAAAGTTTTCTTTTAGCCATAGTAATGCTCTTTGGCATTGCCATGCAGGGACAT
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TCCTTGCGAATGGGACAAACAGCAGTATCTGACAAGATTTCCATCGATTACGTTTATCGG
CAAGGAGATGCTGAGAGGGGTATCACATCACAAGAGGAAGGCTCTCCTGCATATTTTTAT
GTAGCTAATCGTGGAATAATGAGGGCTATGCTCTTGAGCAGCAGATGACAGAATACCG
ACAATTTTAGCCTATTCACCCATTGGCCGTTTTGACATGGACAGTATGCCGGACAATCTT
CGCATGTGGCTACAAATTTACGATCAGGAAATAGGCCTGATACTTTCCGGAAAAGCTCAG
CTCAATGAAGAGATATTACGTACCGAGGGCGTACCGGCTGAAGTACATGCTCTGATGGAT
AACGGTCATTTTGCCAACGATCCCATGCGATGGAATCAAGGTTACCCATGGAACAATAAG
GAACCACTGCTTCCTAATGGCAATCATGCCTATACCGGCTGTGTTGCTACTGCTGCAGCA
CAAATCATGCGCTACCATAGCTGGCCGCTTCAAGGTGAAGGCTCTTTCGATTATCATGCA
GGTTCATTAGTTGGCAACTGGTCCGGCACATTTGGTGAAATGTACGACTGGATCAATATG
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AGCGTTTATGTAGTAGGAGCCTTGCGAAACAACCTTTCGCTACAAGCGTTCACTGCAGCTA
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GGCTTCTACAACTAACACTCCTCTCGCCGACTTCGTTGGGTATCGGAGGTGAGGGAATA
GGTTTTTACCATTTATCAAGAGATCATCACCGGTATCGAACC GGCTAAGACTCCCGCTGAA
GCCGGTACAGATGCCTTGCCGATCTTGGCACTGAAAGACATAGAAGCCGAGTATAAAAGT
GAATCCGGATTGAACGTAGGGTATTCGATATATAATACAGGTGAAGAGCAATCAAATCTT
GACCTCGGATACAGATTGAACAAGGCTGACGGAGAAGTCATAGAGGTGAAAACCTTCATCT
ATCAATATCTCTTGGTACGGATACGGAGAGCATCCCGAGAGTTTCTCATTGGCACCTAAT
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SEQ ID NO:2

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